

# CRF Errors Corrected by the STIC Systems Branch

Serial Number: 09/228516A

CRF Processing Date: 5/9/2002

Edited by: Ar

Verified by: Ar (STIC staff)

#11

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: inserted hard return after Seq. 2

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



1600

## RAW SEQUENCE LISTING

DATE: 05/09/2002

PATENT APPLICATION: US/09/778,516A

TIME: 12:36:48

Input Set : A:\pto.amc.txt

Output Set: N:\CRF3\05092002\I778516A.raw

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5      Lo, Ming Ching
6      Liao, Pei-Ru
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11 <130> FILE REFERENCE: 12875-002001
13 <140> CURRENT APPLICATION NUMBER: 09/778,516A
14 <141> CURRENT FILING DATE: 2001-02-07
16 <150> PRIOR APPLICATION NUMBER: TW 89110235
17 <151> PRIOR FILING DATE: 2000-05-26
19 <160> NUMBER OF SEQ ID NOS: 6
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24 <211> LENGTH: 8115
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Output Set: N:\CRF3\05092002\I778516A.raw

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VERIFICATION SUMMARY

DATE: 05/09/2002

PATENT APPLICATION: US/09/778,516A

TIME: 12:36:49

Input Set : A:\pto.amc.txt

Output Set: N:\CRF3\05092002\I778516A.raw



1636

## RAW SEQUENCE LISTING

DATE: 05/09/2002

PATENT APPLICATION: US/09/778,516A

TIME: 11:48:42

Input Set : A:\12875-002001.txt

Output Set: N:\CRF3\05092002\I778516A.raw

Does Not Comply  
Corrected Diskette Needed

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5       Lo, Ming Ching  
6       Liau, Pei-Ru  
8 <120> TITLE OF INVENTION: LAC SHUTTLE VECTORS  
11 <130> FILE REFERENCE: 12875-002001  
13 <140> CURRENT APPLICATION NUMBER: 09/778,516A  
14 <141> CURRENT FILING DATE: 2001-02-07  
16 <150> PRIOR APPLICATION NUMBER: TW 89110235  
17 <151> PRIOR FILING DATE: 2000-05-26  
19 <160> NUMBER OF SEQ ID NOS: 6  
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0

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P.4

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168 <213> ORGANISM: Lactobacillus plantarum  
170 <400> SEQUENCE: 2

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## RAW SEQUENCE LISTING

DATE: 05/09/2002

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TIME: 11:48:42

Input Set : A:\12875-002001.txt

Output Set: N:\CRF3\05092002\I778516A.raw

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297 cctggccctg ccaaaagaac tgaccgatta ccgtactat ggtctgggac ctaatgagag 7620
298 ctaccagac cgcttggaag gtaattacct gggcatctac caggagcg taaaaaagaa 7680
299 ctttagccca tatcgtccgc aggaacggg caaccggagc aagggtcgct ggtaccagct 7740
300 ctttgatgaa aagggcggct tggaatttac ggccaatggg gcagacttga acttgtctgc 7800
301 ttgccatat tctgccgccc aaattgaagc agcggaccac gcttttgaac tgactaacia 7860
302 ttacacttgg gttagagcct taagcgccca gatgggggtc gccggggatg actcctgggg 7920
303 gcagaaggtc caccgggaat tctgcctgga tgcataaaaa gcccgccagc ttcgcctggt 7980
304 gattcagccc cttttactaa aataaatgct acaattgact taacaggatg aaattttagt 8040
305 aaaagcaaag cgagtgagga agatggcaac gatcagagaa gtgccaaaggc agccggcgctg 8100
E--> 306 tcgctagcga cggtc 8115<210> 3
307 <211> LENGTH: 30
308 <212> TYPE: DNA
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
312 <223> OTHER INFORMATION: oligonucleotide for PCR
W--> 314 <210> SEQ ID NO:
E--> 314 <400> SEQUENCE: 3
315 aagctcatga ttggcagcca gtctccgggc 30
E--> 317 <210> SEQ ID NO: 4

```

insert  
hard  
return

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/778,516A

DATE: 05/09/2002  
TIME: 11:48:43

Input Set : A:\12875-002001.txt  
Output Set: N:\CRF3\05092002\I778516A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:2; Line(s) 306

Skipped Sequences(NEW RULES):

Sequence(s)\_\_\_missing. If intentional, please use the following format for each skipped sequence.

<210> sequence id number

<400> sequence id number

000

Seq#:3